

GenCore version 5.1.6  
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## SUMMARIES

OM nucleic - protein search, using frame\_plus.n2p model

Run on: August 22, 2003, 14:07:07 ; Search time 104 Seconds

(without alignments)  
4740.430 Million cell updates/sec

Title: US-09-745-506-74

Perfect score: 506  
Sequence: 1 GTGATTTGTTATCTGTGCTCTCTCTTTACTTAACATTCAA 1553

Scoring table:

OLIGO  
Xgapop 60.0 , Xgapext 60.0  
Ygapop 60.0 , Ygapext 60.0  
Delop 6.0 , Delext 7.0

Searched: 1107863 seqs, 158726573 residues

Word size: 1

Total number of hits satisfying chosen parameters: 2062474

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Command line parameters:

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-LIST=45 -DOCALLIGN=200 -THR\_SCORE=quality -THR\_MIN=1 -ALIGN=15 -MODE=LOCAL  
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-NO\_MAP -LANG=OTHER -NEG\_SCORES=0 -WAIT -DSPLICE=100 -LONGIDOG  
-DEV\_TIMEOUT=120 -WARN\_TIMEOUT=30 -THREDS=1 -XGAPOP=60 -XGAPEXT=60 -FGAPOP=6  
-FGAPEXT=7 -YGAPOP=60 -YGAPEXT=60 -DELOP=6 -DELEXT=7

Database :

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Result No.	Score	Query Match	Length	DB	ID	Description
1	369	72.9	383	22	AAH88085	Human Immune/haema
2	350	69.2	350	22	AAH81361	Human AFP protein
3	350	69.2	350	22	AAH94573	Human protein sequ
4	315	62.3	377	22	AAH27744	Human full-length
5	268	53.0	351	22	AAH60635	Human gene express
6	211	41.7	247	23	ABH08182	Human protein kina
7	102	20.2	110	22	ABG20985	Human liver peptid
8	68	13.4	68	22	ABH32385	Peptide #5036 enco
9	68	13.4	68	22	ABH37657	Peptide #5173 enco
10	68	13.4	68	22	AAH58285	Human brain expres
11	68	13.4	68	22	AAH18609	Peptide #5043 enco
12	68	13.4	68	22	AAH06178	Peptide #4860 enco
13	49	9.7	79	22	AAH21467	Human novel foetal
14	43	8.5	146	22	AAH27916	Human contig polyp
15	38	7.5	70	22	ABG20982	Human human diagno
16	30	5.9	58	22	AAH90790	Human immune/haema
17	25	4.9	74	22	ABG20984	Human human diagno
18	22	4.4	22	22	ABG20983	Novel human diagno
19	15	3.0	15	23	ABH08183	Human protein kina
20	10	2.0	360	22	AAH82528	S. epidermidis ope
21	10	2.0	367	23	ABH38833	Staphylococcus epi
22	9	1.8	304	23	ABH66109	Blifidobacterium lo
23	8	1.6	10	22	AAH95515	Human complementar
24	8	1.6	52	22	AAH65449	Propionibacterium
25	8	1.6	64	22	AAH21906	Human cardiovascular
26	8	1.6	96	23	ABH70062	Human prey protein
27	8	1.6	108	22	AAH66453	Propionibacterium
28	8	1.6	108	22	AAH92483	C glutamic acid prote
29	8	1.6	161	22	AAH63960	Propionibacterium
30	8	1.6	179	12	AAH10310	Ovary tissue trans
31	8	1.6	218	22	AAH47094	Propionibacterium
32	8	1.6	299	22	AAH24644	Human olfactory re
33	8	1.6	299	22	AAH71682	Human olfactory re
34	8	1.6	299	23	ABH76794	Human G-protein co
35	8	1.6	299	23	AAH95727	Human G-protein co
36	8	1.6	299	23	AAH85264	G-coupled olfactor
37	8	1.6	345	23	AAH13351	Human trypsin protei
38	8	1.6	355	23	ABH91505	Purine/pyrimidine
39	8	1.6	355	24	ABH99273	Orthosomycin biosy
40	8	1.6	355	24	ABH76704	Streptomyces Virid
41	8	1.6	367	19	AAH69999	Rodent chemokine r
42	8	1.6	368	19	AAH54371	Human IP-10/Mig re
43	8	1.6	368	21	AAH90614	Human G protein-co
44	8	1.6	368	21	AAH90614	Human G protein-co
45	8	1.6	368	21	AAH90614	Human mutant G pro

## ALIGNMENTS

RESULT 1  
ID AAH88085 standard; Protein: 383 AA.

AC AAH88085;  
DT 07-NOV-2001 (first entry)

DE Human immune/haematopoietic antigen SEO ID NO:15678.

KW Human; Immune; haematopoietic; Immune/haematopoietic antigen; cancer;  
KW cytostatic; gene therapy; vaccine; metastasis.

OS Homo sapiens.

XX WO200157182-A2.

XX PD 09-AUG-2001.

XX 17-JAN-2001; 2001WO-US01354.  
PF  
PR 31-JAN-2000; 2000US-0179065.  
PR 04-FEB-2000; 2000US-0180628.  
PR 24-FEB-2000; 2000US-0184668.  
PR 02-MAR-2000; 2000US-0186350.  
PR 16-MAR-2000; 2000US-0189874.  
PR 17-MAR-2000; 2000US-0190076.  
PR 18-APR-2000; 2000US-0198123.  
PR 09-MAY-2000; 2000US-0205515.  
PR 07-JUN-2000; 2000US-0209467.  
PR 28-JUN-2000; 2000US-0214886.  
PR 30-JUN-2000; 2000US-0215135.  
PR 07-JUL-2000; 2000US-0216647.  
PR 07-JUL-2000; 2000US-0216880.  
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PR 17-NOV-2000; 2000US-0249265.  
PR 17-NOV-2000; 2000US-0249297.  
PR 17-NOV-2000; 2000US-0249299.  
PR 17-NOV-2000; 2000US-0249300.  
PR 01-DEC-2000; 2000US-0250160.  
PR 01-DEC-2000; 2000US-0250391.  
PR 05-DEC-2000; 2000US-0251030.  
PR 05-DEC-2000; 2000US-0251988.  
PR 06-DEC-2000; 2000US-0256719.  
PR 08-DEC-2000; 2000US-0251856.  
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PR 08-DEC-2000; 2000US-0251889.  
PR 08-DEC-2000; 2000US-0251990.  
PR 11-DEC-2000; 2000US-0254097.  
PR 05-JAN-2001; 2001US-0259676.  
  
(HUMA-) HUMAN GENOME SCI INC.  
PA  
XX  
XX  
PI  
XX  
XX  
DR  
Rosen CA, Barash SC, Ruben SM;  
WPI: 2001-483426/52.  
N-PSDB: AAK60866.

xx nucleic acids encoding human immune/hematopoietic antigen polypeptides,  
 PT useful for preventing, diagnosing and/or treating cancers and  
 PT metastasis -  
 xx  
 xx  
 xx  
 xx  
 xx  
 Claim 11, SEQ ID NO 15678; 3071pp + Sequence Listing; English.  
 CC  
 CC AAK54951 to AAK64702 encode the human immune/hematopoietic antigen (I)  
 CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic  
 CC activity, and can be used in gene therapy and vaccine production. (I)  
 CC proteins and polynucleotides may be used in the prevention, diagnosis and  
 CC treatment of diseases associated with inappropriate (I) expression. For  
 CC example, they may be used to treat disorders associated with decreased  
 CC expression by rectifying mutations or deletions in a patient's genome  
 CC that affect the activity of (I) by expressing inactive proteins or to  
 CC supplement the patients own production of (I). Additionally, (I)  
 CC polynucleotides may be used to produce the secreted (I), by inserting  
 CC the nucleic acids into a host cell and culturing the cell to express the  
 CC protein. (I) proteins and polynucleotides may be used to prevent,  
 CC diagnose and treat immune/hematopoietic-related diseases, especially  
 CC cancers and cancer metastases of hematopoietic-derived cells. AAK64703  
 CC to AAK87694 represent human immune/hematopoietic antigen genomic  
 CC sequences from the present invention. AAK54942 to AAK54950 and AAM82169  
 CC represent sequences used in the exemplification of the present invention

XX	Sequence	383 AA;
Alignment Scores:		
Pred. No.:	0	Length: 383
Score:	369.00	Matches: 369
Percent Similarity:	100.00%	Conservative: 0
Best Local Similarity:	100.00%	Mismatches: 0
Query Match:	72.92%	Indels: 0
DB:	22	Gaps: 0

QY	188	GTCCACGACGACGTCCGGTTTGATGATCCCGATCTGCATATCTCCGCTCCGTATG	24
Db	15	ValProthThrValAlaPheValAspSerLeuIleCysAsnSerSerAlaSerPheMet	34
QY	248	GATTTGAGGCTCTCCCTTCTCCATGATGACTTTGGATCCCTCTCGTTTGGAGAGT	307
Db	35	AspLeuLysAlaLeuLeuSerSerLeuAsnAspPheAlaSerLeuSerPheAlaGlnSer	54
QY	308	TGGGACAAATGTGGATTACTGGTGGACCAAGCCACACATACTGTAAATACACTCTTC	367
Db	55	TirPAspAsnValaGluLeuLeuValaGluProSerProProHisThrValaAsnThrLeuPhe	74
QY	368	CTPACCAATGACCTGACTGAGGAAGGATGAGGAGGAGGCTGCCAAAGAAAGACAGACTC	427
Db	75	LeuThrAsnAspLeuThrGluGluValaMetGluGluValaLeuGlnLysLysAlaAspLeu	94
QY	428	ATTCTCTCCATCACCATCCGGCTATCTCTCCACCAATGAAGCGCATACCTGGAGACAATAG	487
Db	95	IleLeuSerTyrHisIleProIlePheArgPrometLysArgGlyIleThrTirPAsnThrTirP	114
QY	488	AAGGAGCGCTGGTGATCCGGGCTGTGGAGAACAGAGTCGGTATCTCTCTCATACA	547
Db	115	LysGluArgLeuValaIleArgAlaLeuGluGluAsnArgValaGlyIleTyrSerProHisThr	134
QY	548	GCTCATGATGCTCCGGCCCGACGGCGTCACACATCGTTGGCGTAAAGGGCTTGAGACTGT	607
Db	135	AlaTyrAspAlaIleAlaProGlnGlyValaAsnAsnThrPheAlaLysGlyLeuGlyAlaCys	154
QY	608	ACCTCCAGGCGCCATACATCTTCCCAAAGCTCCCACTACCTCAGAGAGAAACACCGCA	667
Db	155	ThrSerArgProIleHisProSerIleLysAlaIleProAsnTyrProThrGluGlyAsnHisArg	174
QY	668	GTAATTAATCAAGCTTAATACACCCAAAGACCTGGACCAAGATGATGTCTGACGTGAAGA	727
Db	175	ValaGluPheAsnValaAsnTyrThrGlnAspLeuAspLysValaMetSerAlaValaLysGly	194

QY	728	ATTGACGGTGTGTTGTGCACCTGTTTGTGTGTGACGTGGTAATGAGAAACAAACACGG	787
Db	195	TTleaspglyValSerValThrSerPheSerIlaaIgtHrGlyaspInGInGlnThrArg	214
QY	788	ATTAAATCGAATTTGACTACAGACGGCTGTATGACGGGGAGAGATTTTCTTCCCGAAC	847
Db	215	TTleasnleuasnCysThrGlnlyalaIeuleuGlnValValAspPheIeuSerIlyGaspn	234
QY	848	AAACAACTTTATTCAGAACACGGAAATTTGTCTACGTGAGAAAGCCTTTGCTCAATACT	907
Db	235	LTyGlnIleuTyGlnLysThrGlnLlIleuSerIleuGlnLysProIleuIleuInIshTr	254
QY	908	GGAAATGGACGGCTATGACACACTGGATGAAATGTCTGCTCCGACACCATATTTGATGGA	967
Db	255	GIyMetGlyArgIeuCysThrIeuaAspIuSerValSerIleuAlaIthMetIleAspArg	274
QY	968	ATTAAAGACACCCAAAACATTCATTCATTTGGCTAGGCCCTTGGGGTGGAGAAACCTTA	1027
Db	275	TTleuysaThIshIeulysIeuSerInIshIleArgIeulAlaIeuGlyValGlyArgThIeu	294
QY	1028	GAGCTCTCAAGCTCAAAAGTCGTGGCCCTGTGATCTGATCTGGGAGACACGCTTGCAGAGT	1087
Db	295	GIuBerGlnValLysValValAlaIaIeucysAlaGlySerGlySerValIleuGlnGly	314
QY	1088	GTTGAGCGTGAACCTTTACCTGCACAGGTGAGATGTCCCATCATGATACTTGGATGCTGCT	1147
Db	315	ValGIuaIlaAspIeuTyIeuThrGlyGluMetSerInIshAspThrIeuaAspIlaIa	334
QY	1148	TGCCAAGGAATAATTCATCTCTGTGTAACAACGAAACACTGAAACGAGGCTTCTTCT	1207
Db	335	SerIInGlyIleAsnValIleIeuCysGlnIshSerAsnThrGlnAlaGlyPheIeuSer	354
QY	1208	GACCTTGCAGATGACGTGGAATTCACACTGGAGAGATAAGATAAATATTATTCATCAGAG	1267
Db	355	AspIeuAspArgMetIleuAspSerInIshIeuGlnAsnLysIleAsnIleIleIeuSerGlu	374
QY	1268	ACTGACAGGACCCCTTTCAGGTGGA 1294	
Db	375	ThrsAspArgAspProIeuGlnValVal 383	

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RESULT 2
AAG81361
ID AAG81361 standard; Protein: 350 AA.
XX
AC AAG81361;
XX
DT 10-SEP-2001 (first entry)
XX
DE Human AFP protein sequence SEQ ID NO:240.
XX
KW Human: secreted protein; secretion; bacterial cell; fungal cell;
KW eukaryotic cell; fusion protein; maltose binding protein;
KW immunoglobulin constant region; polynistidine tag.
XX
OS Homo sapiens.
XX
PN WO200129221-A2.
XX
PD 26-APR-2001.
XX
PF 20-OCT-2000; 2000WO-US29052.
XX
PR 20-OCT-1999; 99US-0160712.
XX
PA (ZYMO ) ZYMOGENETICS INC.
XX
PI Conklin DC, Yee DP;
XX
WP1: 2001-300340/31.
DR N-PSDB; AAH52212.
XX
PT Isolated polypeptide for directing secretion of proteins of interest
from a host cell including, e.g. bacteria, includes contiguous amino

```

PT acid residues of polypeptide with specified amino acids -  
XX Claim 1; Page 424-425; 617pp; English.  
XX AAH52093 to AAH52303 encode the human secreted proteins given in AAH52124  
CC to AAH81453. The secreted proteins can be used for directing the  
CC secretion of proteins of interest from a host cell including bacteria,  
CC fungal cells, and cultured higher eukaryotic cells. The present invention  
CC also describes fusion proteins, where a secreted protein of the invention  
CC is operably linked via a peptide bond or peptide linker to a second  
CC protein selected from the group consisting of maltose binding protein,  
CC an immunoglobulin constant region, a polystyrene tag and a peptide  
CC given in AAH81453.  
XX  
SQ Sequence 350 AA;  
  
Alignment Scores:  
Pred. No.: 0 Length: 350  
Score: 350.00 Matches: 350  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 69.17% Indels: 0  
DB: 22 Gaps: 0  
  
US-09-745-506-74 (1-1553) x AAH81361 (1-350)  
  
QY 245 ATGATTTGAAGGCTCTCTTCTTCTTGAATGATTCCTGCTGCTGAG 304  
DB 1 Metaspseudotschleusserleusnasphalaserleuserphenleu 20  
QY 305 AGTGGGACATGTTGGATCTGCTGACCAAGCCACACATCTGTAATCACTC 364  
DB 21 Sertripsasnavalgyleuvalisnproserprohsthrvalasnthreu 40  
QY 365 TTCCGACCAATGACCTGAGTGAAGTGAAGAGAGAGAGAGAGAGAGAGAG 424  
DB 41 Pheulthrasnapleththrgluvalmetgluvalleugllyslasasp 60  
QY 425 CTCATCTCTCTACACCTCCCTATCTCCGACCATGAAGCCATTAACCTGA 484  
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QY 485 TGGAGAGAGCCCTGCTGATCCGGCTCTGGAACAGAGCGGATCTCTCTAT 544  
DB 81 Trpysgluarglevallearglaleuglunsnargvalglyletyserproh 100  
QY 545 ACAGCCTTATGCTGCGCCCGCCAGGCGCTCAACATGTTGGCTAAAGGCTG 604  
DB 101 Thrilatyrasphalalaproglinslyalasnntprleualalysglyleu 120  
QY 605 TGACCTCCAGCCCATCATCTTCCAAAGCTCCCACTACCTACAGAGGAAAC 664  
DB 121 Cysfthserarprolethhisproserlysalproasnlyrprothrglu 140  
QY 665 CGAGTAGATTCACAGCTTAACCTACACCCAGACCTGGACAAAGTCATG 724  
DB 141 Argvalglupheasnvalasnlyrthrghlnasplasnlyvalmetseral 160  
QY 725 GGAATTCAGGCTGTTCTGCTACCTCTTCTGTAGACCTGGAAAGAGAAACA 784  
DB 161 Glylleasphglyvalservalthrserpheserlalaagthrglyasn 180  
QY 785 CGGATTAATCTGAATTTGACTCAGAAAGCTTTGATGACGTTGATTTCTT 844  
DB 181 Argileasleuasncysrthrghlnlysalaleumetglnvalasphleuser 200  
QY 845 AACCAACACTTTATCAGAGAGAGAAATTTCTGCTACGAGAGAGCTTTG 904  
DB 201 Asnysglntleutyrglnlysthrghlnleuserleugllyslproleu 220  
QY 905 ACTGGAATGGAGGCTTATGACACCTGATGAATGTCCTCCGAGCAACATG 964  
DB 221 Thrlymetcglyargleucysrthrleuasphluservalserleualat 240

QY 965 CGAATAAAGACACCTAAACATATTCATATTCGCTTAGCCCTTGGGGGAGAGACC 1024  
DB 241 Argileysarghlsleuylsleuserhlsilearglaleugllyalglyargthr 260  
QY 1025 TTAGAGCTCAAGCAAGCTGCGCCCTGTGTGTGTCTGAGGAGAGGCTTCGAG 1084  
DB 261 Leugluserglnvallysvallvalalaleucysalaglyserglyservalleu 280  
QY 1085 GGTGTGGGCTGACCTTTACCTCCACAGGTGAGATGCCATCATGATCTTGGATGCT 1144  
DB 281 Glyvalglualaasplethyreuthrlyglumetserhlsiaspthrleuasphla 300  
QY 1145 GCTTCCCAAGAAATTAATGTCATCTCTGTGAAACACAGCAAACTGAAAGGCTTTCTT 1204  
DB 301 Alaserglnlylleasnvalilleucysglnhlseserasnthrghlaargglypheu 320  
QY 1205 TCTGACCTCAGAGATGCTGATCTGATCTGAGATTAAGTAATATATTCATCA 1264  
DB 321 Seraspheuasrmetleuaspserrhlsleuglunsnlyllesnilleuser 340  
QY 1265 GAGACTGACAGGAGACCTCTTCAGGTGTA 1294  
DB 341 Glutthraspargaspproleuglnvalval 350  
  
RESULT 3  
AAB94573  
ID AAB94573 standard; Protein: 350 AA.  
XX  
AC AAB94573;  
XX  
DE 26-JUN-2001 (first entry)  
XX  
XX Human protein sequence SEQ ID NO:15360.  
XX  
XX Human; primer: detection; diagnosis; antisense therapy; gene therapy.  
XX  
OS Homo sapiens.  
XX  
PN Ep1074617-A2.  
XX  
PD 07-FEB-2001.  
XX  
PF 28-JUL-2000; 2000EP-0116126.  
XX  
PR 29-JUL-1999; 99JP-0248036.  
XX  
PR 27-AUG-1999; 99JP-0300253.  
PR 11-JAN-2000; 2000JP-0118776.  
PR 02-MAY-2000; 2000JP-0183767.  
PR 09-JUN-2000; 2000JP-0241899.  
XX  
PA (HELI-) HELIX RES INST.  
XX  
PI Ota T, Isogai T, Nishikawa T, Hayashi K, Saito K, Yamamoto J;  
PI Ishii S, Sugiyama T, Wakamatsu A, Nagai K, Otsuki T;  
DR WPI; 2001-318749/34.  
XX  
PT primer sets for synthesizing polynucleotides, particularly the 5602  
PT full-length cDNAs defined in the specification, and for the detection  
PT and/or diagnosis of the abnormality of the proteins encoded by the  
PT full-length cDNAs -  
XX  
XX Claim 8; SEQ ID 15360; 2537bp + CD ROM; English.  
CC  
CC The present invention describes primer sets for synthesizing 5602  
CC full-length cDNAs defined in the specification, where a primer set  
CC comprises: (a) an oligo-dT primer and an oligonucleotide complementary  
CC to the complementary strand of a polynucleotide which comprises one of  
CC the 5602 nucleotide sequences defined in the specification, where the  
CC oligonucleotide comprises at least 15 nucleotides; or (b) a combination  
CC of an oligonucleotide comprising a sequence complementary to the  
CC complementary strand of a polynucleotide which comprises a 5'-end

CC sequence and an oligonucleotide comprising a sequence complementary to a  
 CC polynucleotide which comprises a 3'-end sequence, where the  
 CC oligonucleotide comprises at least 15 nucleotides and the combination of  
 CC the 5'-end sequence/3'-end sequence is selected from those defined in  
 CC the specification. The primer sets can be used in antisense therapy and  
 CC in gene therapy. The primers are useful for synthesizing polynucleotides,  
 CC particularly full-length cDNAs. The primers are also useful for the  
 CC detection and/or diagnosis of the abnormality of the proteins encoded by  
 CC the full-length cDNAs. The primers allow obtaining of the full-length  
 CC cDNAs easily without any specialized methods. AAH03166 to AAH13628 and  
 CC AAH13633 to AAH18742 represent human cDNA sequences; AAB92416 to  
 CC AAB93893 represent human amino acid sequences; and AAH13629 to AAH13632  
 CC represent oligonucleotides, all of which are used in the exemplification  
 CC of the present invention.

XX Sequence 350 AA:

Alignment Scores:

Pred. No.: 0 Length: 350  
 Score: 350.00 Matches: 350  
 Percent Similarity: 100.00% Conservative: 0  
 Best Local Similarity: 100.00% Mismatches: 0  
 Query Match: 69.17% Indels: 0  
 DB: 22 Gaps: 0

US-09-745-506-74 (1-1553) x AAB94573 (1-350)

QY 245 AAGGATTTGAAGGCTCTCTTCTTCTGGAATGACTTTCACCTCTGTTGCTGAG 304  
 DB 1 MetAspLeuIysAlaLeuLeuSerSerLeuAsnAspPheAlaSerLeuSerPheAlaGlu 20  
 QY 305 AGTTGGACAAATGTTGATTACTGATGGAACCAAGCCACACATCTGTAATATACATC 364  
 DB 21 SerTrpAspAsnValIleLeuLeuValGluProSerProProHisThrValAsnThrLeu 40  
 QY 365 TTCCTGACCAATGACTGACTGAGAGATGATGAGAGAGGCTGCAAAAGAGCAGAC 424  
 DB 41 PheLeuThrAsnAspLeuThrGluGluValMetGluIleValLeuGlnIlySAlaAsp 60  
 QY 425 CTCATCTCTCCACATCGGCTATCTCCGACCATGAGAGGCAATACCTGGAGACACA 484  
 DB 61 LeuIleLeuSerThrIleProPheAlaGlyPheAlaGlyIleThrTrpAsnThr 80  
 QY 485 TGGAGAGAGCGCTGGTATCCGGCTGAGAGACAGAGTCCGATCTCTCTCAT 544  
 DB 81 TrpIysGluArgLeuValIleArgAlaLeuGluAsnArgValIleIleIleIleIleIle 100  
 QY 545 ACAGCCTATGATGCTGCGCCCGCCAGGCGTCAACACTGTTGGCTAAAGGCTTGAGCT 604  
 DB 101 ThrAlaTyrAspAlaAlaProGlnGlyValAsnAsnTrpLeuAlaIlySG:ylleuGlyAla 120  
 QY 605 TGTACTCCAGGCGCCATACATCTTCCAAAGCTCCCAACACCTCAAGGAGGAAACAC 664  
 DB 121 CysThrSerArgProIleHisProSerLysAlaProAsnTrpProThrGluGlyAsnHis 140  
 QY 665 CGAGTGAATTCACCTTAACCTACACCAAGCCTGGACAAAGTCATGTTGAGTGA 724  
 DB 141 ArgValGluPheAsnValAsnTrpThrGlnAspLeuAspIysValMetSerAlaValIlyS 160  
 QY 725 GGAATGAGCGGTTTCTGTCTACTTCTTTTCTGTAGAGCTGTAATAGAGAAACA 784  
 DB 161 GlyIleAspGlyValSerValThrSerPheSerAlaArgThrGlnGlnGlnGlnThr 180  
 QY 785 CGGATTAATCTGAATGTACTGACAGGCTTTCAGAGCTTTCAGAGTTCCTCCCG 844  
 DB 181 ArgIleAsnLeuAsnGlyThrGlnIlySAlaLeuMetGlnValValAspPheLeuSerArg 200  
 QY 845 AACCAACAATTTATCAGAGAGCGAAATCTGTCTACTGAGAGAGCTTGTCTTACAT 904  
 DB 201 AsnIysGlnLeuTrpGlnIlySThrGluIleLeuSerLeuGlnIlySProLeuLeuLeuHis 220  
 QY 905 ACTGGAATGGAGCGTTATGACACACTGAGATGAATCTGTCTCCCTGGCAACCATGATGAT 964  
 DB 221 ThrGlyMetGlyArgLeuCysThrLeuAspGluSerValSerLeuAlaThrMetIleAsp 240  
 QY 965 CGAATAAAGACACCTAAACATATCTCATATTCGCTTACCTTGGGGTGGGAGAAC 1024  
 DB 241 ArgIleLysArgHisLeuLysLeuSerHisIleArgLeuAlaLeuGlyValGlyArgThr 260  
 QY 1025 TTGAGAGCTCAAGTCAAAAGTCGGCCCTGTGTGCTGCTGAGAGCAGCTTTCAG 1084  
 DB 261 LeuGluSerGlnValIlySValIleValAlaLeuCysAlaIleGlySerIleSerValLeuGln 280  
 QY 1085 GGTGTGAGCGTACCTTATCTACACAGATGATGTCATGATGATGATGATGATGATGAT 1144  
 DB 281 GlyValGlnAlaAspLeuTrpLeuThrGlyGluMetSerHisHisAspThrLeuAspAla 300  
 QY 1145 GCTTCCCAAGGATTAATGTCTATCTCTGTGGAACACAGCAACACCTGATGATGAT 1204  
 DB 301 AlaSerGlnGlyIleAsnValIleLeuCysGlnHisSerAsnThrGluArgIlePheLeu 320  
 QY 1205 TCTGACCTTGAGATATGCTGATGATTCACATGAGAGATGAATTAATATTCCTATCA 1264  
 DB 321 SerAspLeuArgAspMetLeuAspSerHisLeuGluAsnIlySLeuSer 340  
 QY 1265 GAGACTGACAGGAGGACCTCTTACAGTGCTA 1294  
 DB 341 GluThrAspArgAspProLeuGlnValVal 350

#### RESULT 4

AAU27744  
 ID AAU27744 standard; Protein: 377 AA.

XX AAU27744;

DE 18-DEC-2001 (first entry)

XX Human full-length polypeptide sequence #69.

XX Mammal; human; rhesus monkey; baker's yeast; fission yeast; Norway rat;  
 KW mouse; Chinese hamster; African clawed frog; fruit fly; dog; leukemata;  
 KW cancer; lymphoma; neuroblastoma; autoimmune disorder; cell proliferation;  
 KW nervous system disorder; inflammatory disorder; cell differentiation;  
 KW angiogenesis; stem cell growth factor; activin; inhibin; cartilage; burn;  
 KW genetic disorder; bone regeneration; tendon; ligament; tissue repair;  
 KW cytoskeletal; antiinflammatory; antiarthritic; vulnary; antiinflammatory;  
 KW antibacterial; immunosuppressive; vasotropic; antiparkinsonian;  
 KW neuroprotective; osteopathic; antidiabetic; antisthmatic; antiallergic;  
 KW immunostimulant; analgesic; gene therapy.

XX Homo sapiens.

XX MO200164834-A2.

XX 07-SEP-2001.

XX 26-FEB-2001; 2001WO-US04926.

XX 28-FEB-2000; 2000US-0515126.

XX 18-MAY-2000; 2000US-0577409.

XX 17-JUN-2000; 2000US-0597707.

XX 14-JUL-2000; 2000US-0616807.

XX 19-SEP-2000; 2000US-0664641.

XX (HYSE-) HYSEQ INC.

XX Tang YT, Liu C, Zhou P, Asundi V, Zhang J, Zhao QA, Ren F,

XX Xue AJ, Yang Y, Wehman T, Wang J, Ma F, Wang D, Chen R, Xu C;

XX Dimaac R;

XX MPI: 2001-589862/66.

XX N-PSDB: AAS44644.

PT Novel polypeptides and nucleic acids obtained from cDNA libraries  
 prepared from various human tissues, for diagnosis, treatment of  
 cancer, neurological, inflammatory disorders and for use in arrays for

Pr detection  
 XX Claim 10; SEQ ID No 241; 153bp; English.  
 PS  
 XX Sequences AAU27676-AAU28019 represent full-length polypeptides and  
 CC cortig polypeptides of the invention. The proteins and their associated  
 CC DNA sequences are useful for the treatment, diagnosis and prevention of  
 CC various types of disorder in a mammalian subject such as a human, dog,  
 CC monkey, mouse, hamster or rat. The disorders include cancers such as  
 CC leukemia, lymphoma and neuroblastoma, autoimmune disorders such as  
 CC multiple sclerosis, connective tissue disease, rheumatoid arthritis,  
 CC diabetes mellitus, allergic rhinitis, asthma and eczema, nervous system  
 CC disorders such as Parkinson's disease, Alzheimer's disease, Huntington's  
 CC chorea, amyotrophic lateral sclerosis, spinal muscular atrophy and  
 CC Wernicke disease, inflammatory disorders such as nephritis, Crohn's  
 CC disease, ischemia-reperfusion injury, shock, sepsis and inflammatory  
 CC bowel disease. The sequences exhibit activity relating to angiogenesis,  
 CC cell proliferation, cell differentiation, stem cell growth factor,  
 CC activin or inhibin. Therefore, they can be used to manipulate stem cells  
 CC in culture to give rise to neuroepithelial cells that can be used to  
 CC augment or replace cells damaged by illness, accidental damage or genetic  
 CC disorders. The sequences may also be used for regeneration of bone,  
 CC cartilage, tendons and ligaments and in tissue repair and burn healing.  
 CC Note: Some sequences for this patent did not form part of the printed  
 CC specification, but were obtained in electronic format directly from WIPO  
 CC at ftp.wipo.int/pub/published\_pct\_sequences.  
 CC  
 XX  
 SQ Sequence 377 AA;  
 Alignment Scores:  
 Pred. No.: 3,92e-312 Length: 377  
 Score: 315.00 Matches: 315  
 Percent Similarity: 100.00% Conservative: 0  
 Best Local Similarity: 100.00% Mismatches: 0  
 Query Match: 62.25% Indels: 0  
 DB: 22 Gaps: 0  
 US-09-745-506-74 (1-1553) x AAU27744 (1-377)  
 QY 188 GTCCCAAGACAGATCCGGTGTGTAGATCCCTGATCTGCAATTCCTCCGTTCTTCATG 247  
 DB 9 ValProthrhValArgPheValAspSerLeuIleCysAsnSerSerArgSerPheMet 28  
 QY 248 GATTGGAAGCTCTCTCTCTCTCTCTGATGATGATGATGATGATGATGATGATGATGAT 307  
 DB 29 AspLeuYsaIalLeuLeuSerSerLeuAsnAspPheAlaSerLeuSerPheAlaGluSer 48  
 QY 308 TGGGACATGTGGATTACTGTGAGACCAAGCCACACATACCTGTAATACACTCTTC 367  
 DB 49 TrpAspAsnValGlyLeuLeuValGluProSerProPheHisThrValAsnThrLeuPhe 68  
 QY 368 CTGACCAATGACTGCTGAG 427  
 DB 69 LeuThrAsnAspLeuThrGluGluValMetGluGluValLeuGluGluValAlaAspLeu 88  
 QY 428 ATTCTCTCTACCATCCGCTATCTTCGACCCATGAAGCCATACCTGGAGACATG 487  
 DB 89 IleuSerTyrHisProPheIlePheArgProMetLysArgIleThrTrpAsnThrTrp 108  
 QY 488 AAGAGAGCCCTGGGATCCGGGCTGTGAGAACAGAGAGAGAGAGAGAGAGAGAGAGAGAG 547  
 DB 109 LysGluArgLeuValIleArgAlaLeuGluAsnArgValGlyIleTyrSerProHisThr 128  
 QY 548 GCCTATGATGCTGGCCGCCAGAGGCGTCAACAACAGTGGTGAAGGCTTGGAGCTTGT 607  
 DB 129 AlaTyrAspAlaAlaProGlnGlyValAsnAsnTrpLeuAlaLysGlyLeuGluLysLacys 148  
 QY 608 ACCTGACGCGCCATACATCTCTCCAAAGCTCCCAACTACCTACAGAGGAAACCCAGCA 667  
 DB 149 ThrSerArgProIleHisProSerLysAlaProAsnTyrProThrGluGlyAsnHisArg 168  
 QY 668 GTAGAAATCAACGTTAACTACACCCAGACCTGGAGCAAGCATGTGCGAGTGAAGA 727

DB 169 ValGluPheAsnValAsnTyrThrGlnAspLeuAspLysValMetSerAlaValLysGly 188  
 QY 728 ATTGACGGTGTCTTGTGCTACTTCTTTCTGCTAGACTGGTAATGAGAAACAACAGC 787  
 DB 189 IleAspGlyValSerValThrSerPheSerAlaArgThrGlyAsnGluGluGlnThrArg 208  
 QY 788 ATTAATCTGATTTGATTCACAGAGAGCTTTGATGCGAGGTGTGATTTCTTCCCGAAC 847  
 DB 209 IleAsnLeuAsnCysThrGlnLysAlaLeuMetGlnValValAspPheLeuSerTrpAsn 228  
 QY 848 AAACACCTTTTTCACAAGACGGAATTTCTGCACAGAGAGAGAGAGAGAGAGAGAGAGAG 907  
 DB 229 LysGlnLeuTyrGlnLysThrGluIleLeuSerLeuGluLysProLeuLeuHisThr 248  
 QY 908 GGAATGGAGCGGTATTCACACATGATGATCTGTCTCCCTGGCAACATGATGATGCA 967  
 DB 249 GlyMetGlyArgLeuGlyCysThrLeuAspGlnSerValSerLeuAlaThrMetIleAspArg 268  
 QY 968 ATAAAGACACACCTTAAACATATCTCATATTGCGCTTACGCCCTTGGGGGTGGAGAACCTTA 1027  
 DB 269 IleLysArgHisLeuLysLeuSerHisLeuArgLeuAlaLeuGlyValGlyArgThrLeu 288  
 QY 1028 GAGCTCAAGTCAAGTGTGGCCCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 1087  
 DB 289 GlnSerGlnValLysValValAlaLeuGlySalGlySerGlySerValLeuGlnGly 308  
 QY 1088 GTTGAGCGTCACTTACCTCAGAGTGAATGTCATCATGAT 1132  
 DB 309 ValGluAlaAspLeuTyrLeuThrGlyGluMetSerHisHisAsp 323  
 RESULT 5  
 AAB60663  
 ID AAB60663 standard; Protein; 351 AA.  
 XX  
 AC AAB60663;  
 XX  
 DT 04-MAY-2001 (first entry)  
 XX  
 DE Human gene expression regulatory factor-related protein hnrf3-s.  
 XX  
 KW Human gene expression regulatory factor-related protein; hnrf3-s;  
 KW NGG1-interacting factor; haemopoietic stem cell; preparation;  
 KW detection.  
 XX  
 OS Homo sapiens.  
 XX  
 PN CN1272543-A.  
 PD 08-NOV-2000.  
 XX  
 PF 11-APR-2000; 2000CN-0115369.  
 XX  
 PR 11-APR-2000; 2000CN-0115369.  
 XX  
 PA (NANF-) NANFANG RES CENT STATE HUMAN GENE GROUP.  
 XX  
 LI Li N, Xiao H, Liu F;  
 PI WPI; 2001-183596/19.  
 DR N-PSDB; AAF59945.  
 XX  
 PT Human gene expression regulatory factor related protein and its coded  
 PT sequence -  
 XX  
 Claim 4; Page 19-20; 20pp; Chinese.  
 CC The invention relates to a novel human gene expression regulatory  
 CC factor-related protein, hnrf3-s (NGG1-interacting factor, AAB60663),  
 CC and cDNA encoding it (AAF59945). hnrf3-s is expressed in haemopoietic  
 CC stem cells. The invention also relates to the preparation of hnrf3-s  
 CC proteins and nucleic acids, and the detection of hnrf3-s proteins and  
 CC nucleic acids in a sample. The present sequence represents hnrf3-s.





Db 157 ThrleugluserglnVallyValValAlaLeucysalaglySerGlySerSerValleu 176  
Qy 1082 CAGGGTGTGGAGTGGACCTTACCTCAGAGTGATGTCACATCATGATCTTGGAT 1141  
Db 177 GlnGlyValGlnAlaAspLeuThrGlyGlnMetSerHisHisAspThrLeuASP 196  
Qy 1142 GCTGCTTCCCAAGGAATTAATGTCATCTCTGTGACACACACACACTGACGAGGCTTT 1201  
Db 197 AlaAlaSerGlnGlyIleAsnValIleLeucCysGlnHisSerAsnThrGlnArgGlyPhe 216  
Qy 1202 CTTTCTGACCTTCGAGATATGCTGATTTCTCACTTGGAGATTAAGATTAATATTATCTTA 1261  
Db 217 LeuSerAspLeuArgAspMetLeuAspSerHisLeuGlnAsnLysIleAsnIleLeu 236  
Qy 1262 TCAGAGACTGACAGGAGGACCTCTTCAGTGCTGA 1294  
Db 237 SerGlnThrAspArgAspProLeuGlnValVal 247  
RESULT 7  
ABG20985  
ID ABG20985 standard; Protein; 110 AA.  
AC ABG20985;  
XX  
XX 18-FEB-2002 (first entry)  
DT  
XX  
XX Novel human diagnostic protein #20976.  
DE  
XX Human; chromosome mapping; gene mapping; gene therapy; forensic;  
KW food supplement; medical imaging; diagnostic; genetic disorder.  
XX  
XX Homo sapiens.  
OS  
XX WO200175067-A2.  
PN  
XX 11-OCT-2001.  
PD  
XX 30-MAR-2001; 2001WO-US08631.  
PF  
XX 31-MAR-2000; 2000US-0540217.  
PR 23-AUG-2000; 2000US-0649167.  
XX  
XX (HYSE-) HYSEQ INC.  
PA  
XX Drmanac RT, Liu C, Tang YT;  
PI WPI; 2001-639362/73.  
DR N-PSDB; AAS85172.  
XX  
XX Claim 20; SEQ ID No 51344; 103pp; English.  
PS  
XX The invention relates to isolated polynucleotide (I) and  
CC polypeptide (II) sequences. (I) is useful as hybridisation probes,  
CC polymerase chain reaction (PCR) primers, oligomers, and for chromosome  
CC and gene mapping, and in recombinant production of (II). The  
CC polynucleotides are also used in diagnostics as expressed sequence tags  
CC for identifying expressed genes. (I) is useful in gene therapy techniques  
CC to restore normal activity of (II) or to treat disease states involving  
CC (II). (II) is useful for generating antibodies against it, detecting or  
CC quantitating a polypeptide in tissue, as molecular weight markers and as  
CC a food supplement. (II) and its binding partners are useful in medical  
CC imaging of sites expressing (II). (I) and (II) are useful for treating  
CC disorders involving aberrant protein expression or biological activity.  
CC The polypeptide and polynucleotide sequences have applications in  
CC diagnostics, forensics, gene mapping, identification of mutations  
CC and responsible for genetic disorders or other traits to assess biodiversity  
CC and to produce other types of data and products dependent on DNA and  
CC amino acid sequences. ABG00010-ABG30377 represent novel human

CC diagnostic amino acid sequences of the invention.  
CC Note: The sequence data for this patent did not appear in the printed  
CC specification, but was obtained in electronic format directly from WIPO  
CC at ftp.wipo.int/pub/published\_pcl\_sequences.  
XX  
SQ Sequence 110 AA;  
Alignment Scores:  
Pred. No.: 1e-94 Length: 110  
Score: 102.00 Matches: 102  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 20.16% Indels: 0  
DB: Gaps: 0  
US-09-745-506-74 (1-1553) x ABG20985 (1-110)  
Qy 764 ACTGGAATGAGAAACAACACGGAATTATGTGAATTGTACTGAAAGCCTTTCATGACG 823  
Db 2 ThrGlyAsnGlnGlnGlnThrArgIleAsnLeuAsnCysThrGlnLysAlaLeuMetGln 21  
Qy 824 GTGGTAGATTTCTTCCCGGACAAACAATTATCAGAGACGAAATTTCTGCACTG 883  
Db 22 ValValAspPheLeuSerArgAsnLysGlnLeuThrGlnLysThrGlnIleLeuSerLeu 41  
Qy 884 GAGAAACCTTTGCTTTTACATATCTGGAATGGGACGCTTATGCACACTGGATGAATCTGTC 943  
Db 42 GlnLysProLeuLeuLeuHisThrGlyMetGlyArgLeuCysThrLeuAspGlnSerVal 61  
Qy 944 TCCCTGGGACCAATGATGATGCAATTAATAAGACACCTTAATGTCATATTCGCTTA 1003  
Db 62 SerLeuAlaThrMetIleAspArgIleLysArgHisLeuLysLeuSerHisIleArgLeu 81  
Qy 1004 GCCCTTGGGCTGGGAGAACCTTGAAGTCAAGTCAAAAGCTGGCCCTGTGCTGTG 1063  
Db 82 AlaLeuGlyValGlyArgThrLeuGlnSerGlnValValAlaLeuCysAlaGly 101  
Qy 1064 TCTGGG 1069  
Db 102 SerGly 103  
RESULT 8  
ABG52473  
ID ABG52473 standard; Peptide; 68 AA.  
XX  
XX ABG52473;  
AC  
XX 25-FEB-2003 (first entry)  
DT  
XX Human liver peptide, SEQ ID No 31121.  
DE  
XX Human; liver; cirrhosis; hyperlipoproteinaemia; hyperlipidaemia;  
KW hypercholesterolaemia; coronary heart disease.  
XX  
XX Homo sapiens.  
OS  
XX WO200157273-A2.  
PN  
XX 09-AUG-2001.  
PD  
XX 30-JAN-2001; 2001WO-US00664.  
PF  
XX 04-FEB-2000; 2000US-0180312.  
PR 26-MAY-2000; 2000US-0207456.  
PR 30-JUN-2000; 2000US-0608408.  
PR 03-AUG-2000; 2000US-0632366.  
PR 21-SEP-2000; 2000US-0234687.  
PR 27-SEP-2000; 2000US-0236359.  
PR 04-OCT-2000; 2000GB-0024263.  
PA (MOLE-) MOLECULAR DYNAMICS INC.  
XX Penn SG, Hanzel DK, Chen W, Rank DR;  
PI



XX WPI; 2001-488898/53.  
XX Human genome-derived single exon nucleic acid probes useful for  
PT analysing gene expression in human adult liver.  
XX  
XX Claim 27; SEQ ID NO 31121; 658bp; English.  
XX  
CC The invention relates to a single exon nucleic acid probe (SENP) (I) for  
CC measuring human gene expression in a sample derived from human adult  
CC liver, comprising one of 13109 defined nucleotide sequences given in the  
CC specification (or complements/fragments). The probe hybridises at high  
CC stringency to a nucleic acid molecule expressed in the human adult  
CC liver. (I) may be used for predicting, measuring and displaying gene  
CC expression in samples derived from human adult liver. The genes  
CC identified may be involved in genetic liver diseases such as cirrhosis,  
CC hypeliproteinaemia, hyperlipidaemia and hypercholesterolaemia which  
CC is associated with coronary heart disease. AbG47348-ABG59930 represent  
CC human liver single exon encoded peptides of the invention.  
CC Note: The sequence information for this patent does not appear in the  
CC printed specification but was obtained in electronic format directly  
CC from WIPO at [http://wipo.int/pub/published\\_pct\\_sequences](http://wipo.int/pub/published_pct_sequences).  
XX  
SQ Sequence 68 AA;  
Alignment Scores:  
Pred. No.: 5 29e-60 Length: 68  
Score: 68.00 Matches: 68  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 13.44% Indels: 0  
DB: 22 Gaps: 0  
US-09-745-506-74 (1-1553) x ABG52473 (1-68)  
QY 395 ATGGAGAGGTGCTGCAAAAGAGCAGACCTATTCTCTCCATCCGCTATCTTC 454  
DB 1 MetGluGluValLeuGlnLysAlaAspLeuIleuSerYrHisProProlIePhe 20  
QY 455 CGACCATGAAGGCGATTAACCTGGAACATGGAAGAGCGCTGGTGCATCCGGCTCTG 514  
DB 21 ArgProMetLysArgIleThrTrpAsnThrTrpLysGluIuArgLeuValIleArgAlaLeu 40  
QY 515 GAGACAGAGTCGGTATCTACTCTCCATACAGCCTATGATGCTGGCCGAGGGGCTC 574  
DB 41 GluAsnArgValGlyIleTyrSerProHisThrAlaIleArgAlaProGlnGlyVal 60  
QY 575 AACCACTGGTGGCTAAAGGGCTT 598  
DB 61 AsnAsnTrpLeuAlaLysGlyLeu 68  
RESULT 9  
ABB32385  
ID ABB32385 standard; Peptide: 68 AA.  
XX  
XX ABB32385;  
DT 01-FEB-2002 (first entry)  
XX  
DE Peptide #5036 encoded by breast cell single exon nucleic acid probe.  
XX  
XX Human: microarray; single exon probe; gene expression; breast;  
KM disease; cancer.  
XX  
XX Homo sapiens.  
XX  
XX WO200157271-A2.  
XX  
XX PD 09-AUG-2001.  
XX  
XX 30-JAN-2001; 2001MO-US00662.  
XX  
XX 04-FEB-2000; 2000US-0180312.  
XX

PR 26-MAY-2000; 2000US-0207456.  
PR 30-JUN-2000; 2000US-0608408.  
PR 03-AUG-2000; 2000US-0632366.  
PR 21-SEP-2000; 2000US-0234687.  
PR 27-SEP-2000; 2000US-0236359.  
PR 04-OCT-2000; 2000GB-0024263.  
XX  
XX (MOLE-) MOLECULAR DYNAMICS INC.  
XX  
XX Penn SG, Hanzel DK, Chen W, Rank DR;  
PI WPI; 2001-486933/54.  
XX  
XX New spatially-addressable set of single exon nucleic acid probes,  
PT useful for measuring gene expression in sample derived from human  
PT breast, comprises number of single exon nucleic acid probes -  
XX  
XX Claim 27; SEQ ID NO 15353; 327pp + sequence listing; English.  
XX  
XX The invention relates to a spatially-addressable set of single exon  
CC nucleic acid probes for measuring gene expression in a sample derived  
CC from human breast and BT 474 cells. The method involves contacting  
CC the probes with a collection of detectably labelled nucleic acids  
CC derived from mRNA of human breast, and then measuring the label  
CC bound to each probe of the microarray. The probes are useful for  
CC verifying the expression of regions of genomic DNA predicted to  
CC encode proteins. They are useful for gene discovery, and for  
CC determining predisposition and/or prognosing breast disease. Gene  
CC expression analysis is useful for assessing the toxicity of chemical  
CC agents on cells. The microarray of this invention presents a far greater  
CC diversity of probes for measuring gene expression, with far less bias  
CC than expressed sequence tag microarrays. The method is suitable for  
CC rapid production of functional information from genomic sequence. The  
CC present sequence is a peptide encoded by a single exon nucleic acid  
CC probe of the invention.  
CC Note: The sequence data for this patent did not form part of the  
CC printed specification, but was obtained in electronic format directly  
CC from WIPO at [http://wipo.int/pub/published\\_pct\\_sequences](http://wipo.int/pub/published_pct_sequences).  
XX  
SQ Sequence 68 AA;  
Alignment Scores:  
Pred. No.: 5 29e-60 Length: 68  
Score: 68.00 Matches: 68  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 13.44% Indels: 0  
DB: 22 Gaps: 0  
US-09-745-506-74 (1-1553) x ABB32385 (1-68)  
QY 395 ATGGAGAGGTGCTGCAAAAGAGCAGACCTATTCTCTCCATCCGCTATCTTC 454  
DB 1 MetGluGluValLeuGlnLysAlaAspLeuIleuSerYrHisProProlIePhe 20  
QY 455 CGACCATGAAGGCGATTAACCTGGAACATGGAAGAGCGCTGGTGCATCCGGCTCTG 514  
DB 21 ArgProMetLysArgIleThrTrpAsnThrTrpLysGluIuArgLeuValIleArgAlaLeu 40  
QY 515 GAGACAGAGTCGGTATCTACTCTCCATACAGCCTATGATGCTGGCCGAGGGGCTC 574  
DB 41 GluAsnArgValGlyIleTyrSerProHisThrAlaIleArgAlaProGlnGlyVal 60  
QY 575 AACCACTGGTGGCTAAAGGGCTT 598  
DB 61 AsnAsnTrpLeuAlaLysGlyLeu 68  
RESULT 10  
ABB37667  
ID ABB37667 standard; Peptide: 68 AA.  
XX  
XX ABB37667;  
XX

DT 04-FEB-2002 (first entry)  
XX  
XX Peptide #5173 encoded by human foetal liver single exon probe.  
XX  
XX Human; foetal liver; gene expression; single exon nucleic acid probe.  
XX  
OS Homo sapiens.  
XX  
XX MO200157277-A2.  
XX  
XX 09-AUG-2001.  
XX  
XX 30-JAN-2001; 2001WO-US00669.  
XX  
XX 04-FEB-2000; 2000US-0180312.  
XX 26-MAY-2000; 2000US-0207456.  
XX 30-JUN-2000; 2000US-0608408.  
XX 03-AUG-2000; 2000US-0632366.  
XX 21-SEP-2000; 2000US-0234687.  
XX 27-SEP-2000; 2000US-0236359.  
XX 04-OCT-2000; 2000GB-0024263.  
XX  
XX (MOLE-) MOLECULAR DYNAMICS INC.  
XX  
XX Penn SG, Hanzel DK, Chen W, Rank DR;  
XX  
XX WPI; 2001-483447/52.  
XX  
XX Human genome-derived single exon nucleic acid probes useful for  
PT analyzing gene expression in human fetal liver -  
PT  
XX  
XX Claim 27; SEQ ID NO 30302; 639pp + sequence listing; English.  
XX  
XX The invention relates to a single exon nucleic acid probe for  
CC measuring human gene expression in a sample derived from human foetal  
CC liver. The single exon nucleic acid probes may be used for predicting,  
CC measuring and displaying gene expression in samples derived from human  
CC fetal liver. The present sequence is a peptide encoded by a single exon  
CC nucleic acid probe of the invention.  
CC Note: The sequence data for this patent did not form part of the  
CC printed specification, but was obtained in electronic format directly  
CC from WIPO at ftp.wipo.int/pub/published\_pct\_sequences.  
XX  
XX  
SQ Sequence 68 AA:  
Alignment Scores:  
Pred. No.: 5,29e-60 Length: 68  
Score: 68.00 Matches: 68  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 13.44% Indels: 0  
DB: 22 Gaps: 0  
US-09-745-506-74 (1-1553) x ABB37667 (1-68)  
QY 395 ATGGAGAGGTGCTGCAAGAAGACACCTATTCTCTACCATCGGCTATCTTC 454  
DB 1 MetGUGluValLeuGlnLysLysAlaSplLeuIleuSerTYrHisProProlIePhe 20  
QY 455 CGACCATGAAGCGCATACCTGGAACACATGGAAGAGCGCGTGTATCCGGGCTGTC 514  
DB 21 ArgPrometLysArgIleThrTyrPasnThrTrpLysGluArgLeuValIleArgAlaLeu 40  
QY 515 GAGAACAAGATCGGTATCTACTCTCTCATACAGGCTATGATGCTGCGCCCGAGGCGCTC 574  
DB 41 GluAsnArgValAlGlyIleTyrSerProHisThrAlaTyrAspAlaAlaProGlnGlyVal 60  
QY 575 AACACACTGTTGGCTAAAGGCTT 598  
DB 61 AsnAsnTrpLeuAlaIalysGlyLeu 68  
RESULT 11  
AAM58295

ID AAM58295 standard; Protein: 68 AA.  
XX  
XX  
XX AAM58295;  
XX  
XX 05-NOV-2001 (first entry)  
XX  
XX Human brain expressed single exon probe encoded protein SEQ ID NO: 30400.  
DE  
XX  
XX Human; brain expressed exon; gene expression analysis; probe;  
KW microarray; Alzheimer's disease; multiple sclerosis; schizophrenia;  
KW epilepsy; cancer.  
XX  
XX Homo sapiens.  
XX  
XX MO200157275-A2.  
XX  
XX 09-AUG-2001.  
XX  
XX 30-JAN-2001; 2001WO-US00667.  
XX  
XX 04-FEB-2000; 2000US-0180312.  
XX 26-MAY-2000; 2000US-0207456.  
XX 30-JUN-2000; 2000US-0608408.  
XX 03-AUG-2000; 2000US-0632366.  
XX 21-SEP-2000; 2000US-0234687.  
XX 27-SEP-2000; 2000US-0236359.  
XX 04-OCT-2000; 2000GB-0024263.  
XX  
XX (MOLE-) MOLECULAR DYNAMICS INC.  
XX  
XX Penn SG, Hanzel DK, Chen W, Rank DR;  
XX  
XX WPI; 2001-483446/52.  
XX  
XX Single exon nucleic acid probes for analyzing gene expression in human  
PT brains -  
PT  
XX  
XX Example 4; SEQ ID NO: 30400; 650pp + sequence listing; English.  
XX  
XX The present invention provides a number of single exon nucleic acid  
CC probes which are derived from genomic sequences expressed in the human  
CC brain. They can be used to measure gene expression in brain cell samples,  
CC which may enable the diagnosis and improved treatment of nervous system  
CC diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia,  
CC epilepsy and cancers. The present sequence is a protein encoded by one of  
CC the probes of the invention.  
XX  
XX  
SQ Sequence 68 AA:  
Alignment Scores:  
Pred. No.: 5,29e-60 Length: 68  
Score: 68.00 Matches: 68  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 13.44% Indels: 0  
DB: 22 Gaps: 0  
US-09-745-506-74 (1-1553) x AAM58295 (1-68)  
QY 395 ATGGAGAGGTGCTGCAAGAAGACACCTATTCTCTACCATCGGCTATCTTC 454  
DB 1 MetGUGluValLeuGlnLysLysAlaSplLeuIleuSerTYrHisProProlIePhe 20  
QY 455 CGACCATGAAGCGCATACCTGGAACACATGGAAGAGCGCGTGTATCCGGGCTGTC 514  
DB 21 ArgPrometLysArgIleThrTyrPasnThrTrpLysGluArgLeuValIleArgAlaLeu 40  
QY 515 GAGAACAAGATCGGTATCTACTCTCTCATACAGGCTATGATGCTGCGCCCGAGGCGCTC 574  
DB 41 GluAsnArgValAlGlyIleTyrSerProHisThrAlaTyrAspAlaAlaProGlnGlyVal 60  
QY 575 AACACACTGTTGGCTAAAGGCTT 598  
AAM58295

Db 61 AsnAsnTrpLeuAlaLysGlyLeu 68

RESULT 12  
AAM18609  
ID AAM18609 standard; Protein; 68 AA.  
XX  
AC AAM18609;  
XX  
DT 12-OCT-2001 (first entry)  
XX  
DE Peptide #5043 encoded by probe for measuring cervical gene expression.  
XX  
KW Probe; human; microarray; gene expression; cervical epithelial cell;  
XX  
KW cervical cancer.  
XX  
OS Homo sapiens.  
XX  
PN MO200157278-A2.  
XX  
PD 09-AUG-2001.  
XX  
PF 30-JAN-2001; 2001MO-US00670.  
XX  
PR 04-FEB-2000; 2000US-0180312.  
XX  
PR 26-MAY-2000; 2000US-0207456.  
XX  
PR 30-JUN-2000; 2000US-0608408.  
XX  
PR 03-AUG-2000; 2000US-0632366.  
XX  
PR 21-SEP-2000; 2000US-0234687.  
XX  
PR 27-SEP-2000; 2000US-0236359.  
XX  
PR 04-OCT-2000; 2000GB-0024263.  
XX  
PA (MOLE-) MOLECULAR DYNAMICS INC.  
XX  
PI Penn SG, Hanzel DK, Chen W, Rank DR;  
XX  
DR WPI; 2001-488901/53.  
XX  
PT Human genome-derived single exon nucleic acid probes useful for  
XX  
PS Claim 27; SEQ ID NO 23435; 487pp; English.  
XX  
CC The present invention relates to human single exon nucleic acid probes  
CC (SEN; see AAI10068-AA128459). The present sequence is a peptide encoded  
CC by one such probe. The SENs are derived from human HeLa cells. The SENs  
CC can be used to produce a single exon microarray, which can be used for  
CC measuring human gene expression in a sample derived from human cervical  
CC epithelial cells. By measuring gene expression, the probes are therefore  
CC useful in grading and/or staging of diseases of the cervix, notably  
CC cervical cancer.  
CC Note: The sequence data for this patent did not form part of the printed  
CC specification, but was obtained in electronic format directly from WIPO  
CC at ftp.wipo.int/pub/published\_pct\_sequences.  
XX  
SQ Sequence 68 AA;  
XX

Alignment Scores:  
Pred. No.: 5.29e-60 Length: 68  
Score: 68.00 Matches: 68  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 13.44% Indels: 0  
DB: 22 Gaps: 0

US-09-745-506-74 (1-1553) x AAM18609 (1-68)

OY 395 ATGGAGAGGTCTGCAAAAAGAGGAGACCTATCTCTCCATCCATCCGCTATCTTC 454  
Db 1 MerglUGlVAlleuGlnLysLysAlaAplLeuIlleuSerYrHISpROpOlllePhe 20  
OY 455 CGACCATGAGGCGCTAACCTGGAACACATGAGAGAGCGCTGTGATCCGGGCTCTG 514  
Db 21 AtgPrometLysArgIleThrTrrPasnThrTrpLysGluArgLeuValIleArgAlaLeu 40

OY 515 GAGACAGAGTCGTATCTACTCTCCATACAGACCTATGATGCTGCGCCAGGCGCTC 574  
Db 41 GluAsnArgValGlyIleYrSerProHisThrAlaLysrPspAlaIAPROGInGlyVal 60  
OY 575 AACACTGTGTGGCTAAAGGCGTT 598  
Db 61 AsnAsnTrpLeuAlaLysGlyLeu 68

RESULT 13  
AAM06178  
ID AAM06178 standard; Protein; 68 AA.  
XX  
AC AAM06178;  
XX  
DT 09-OCT-2001 (first entry)  
XX  
DE Peptide #4860 encoded by probe for measuring breast gene expression.  
XX  
KW Probe; human; breast disease; breast cancer; development disorder;  
XX  
KW inflammatory disease; proliferative breast disease; non-carcinoma tumour.  
XX  
OS Homo sapiens.  
XX  
PN MO200157270-A2.  
XX  
PD 09-AUG-2001.  
XX  
PF 29-JAN-2001; 2001MO-US00661.  
XX  
PR 04-FEB-2000; 2000US-0180312.  
XX  
PR 26-MAY-2000; 2000US-0207456.  
XX  
PR 30-JUN-2000; 2000US-0608408.  
XX  
PR 03-AUG-2000; 2000US-0632366.  
XX  
PR 21-SEP-2000; 2000US-0234687.  
XX  
PR 27-SEP-2000; 2000US-0236359.  
XX  
PR 04-OCT-2000; 2000GB-0024263.  
XX  
PA (MOLE-) MOLECULAR DYNAMICS INC.  
XX  
PI Penn SG, Hanzel DK, Chen W, Rank DR;  
XX  
DR WPI; 2001-476286/51.  
XX  
PT Novel single exon nucleic acid probe used to measuring gene expression  
XX  
PS Claim 27; SEQ ID NO 14918; 322pp; English.  
XX  
CC The present invention relates to novel single exon nucleic acid probes  
CC (see AAI00010-AA110067). The present sequence is a peptide encoded by one  
CC such probe. The probes are useful for measuring human gene expression in  
CC a human breast sample, where the probe hybridises at high stringency to a  
CC nucleic acid expressed in the human breast. The probes are useful for  
CC predicting, diagnosing, grading, staging, monitoring and prognosing  
CC diseases of the human breast, particularly those diseases with polygenic  
CC aetiology. The diseases include: breast cancer, disorders of development,  
CC inflammatory diseases of the breast, fibrocystic changes, proliferative  
CC breast disease and non-carcinoma tumours.  
CC Note: The sequence data for this patent did not form part of the printed  
CC specification, but was obtained in electronic format directly from WIPO  
CC at ftp.wipo.int/pub/published\_pct\_sequences.  
XX  
SQ Sequence 68 AA;  
XX

Alignment Scores:  
Pred. No.: 5.29e-60 Length: 68  
Score: 68.00 Matches: 68  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 13.44% Indels: 0  
DB: 22 Gaps: 0

US-09-745-506-74 (1-1553) x AAM06178 (1-68)

OY 395 ATGGAGGAGGTGTCGCAAAAGAGCAGACCTCATTCCTCTACCATCGGCTATCTTC 454  
|||||  
Db 1 MetcUguvAlldGlnLysAlaAspLeuIleuSerTyrHisProIleIle 20  
OY 455 CGACCCATGAGAGCCATTAACCTGGAACACATGGAAGAGAGCCCTGGTATCGGGCTTG 514  
|||||  
Db 21 ArgProMetLysArgIleThrTrpAsnThrTrpLysGluArgLeuValIleArgIle 40  
OY 515 GAGAAACAGTGGGATCTACTCTCCATACAGCCTATGATGCTGGCCGCCAGGCGCTC 574  
|||||  
Db 41 GlusnaArgValIGlyIleTyrSerProHisThrAlaTyrAspAlaAlaProGlnIleVal 60  
OY 575 AACAACTGTGTGGCTAAAGGCTT 598  
|||||  
Db 61 AsnAsnTrpLeuAlaIalysGlyLeu 68

RESULT 14  
AAU21467  
ID AAU21467 standard; Protein; 79 AA.  
XX  
AC AAU21467;  
XX  
DT 18-DEC-2001 (first entry)  
XX  
DE Human novel foetal antigen; SEQ ID NO 1711.  
XX  
KW Human; foetal tissue antigen; antiinflammatory; neuroprotective;  
KW Immunomodulator; cardiovascular; cytosolic; nephrothropic;  
KW Cardiovascular; autoimmune disease; rheumatoid arthritis;  
KW hyperproliferative disorder; breast neoplasm; cancer;  
KW cardiovascular disorder; cardiac arrest; cerebrovascular disorder;  
KW cerebral ischemia; angiogenesis; nervous system disorder;  
KW Alzheimer's disease; infection; ocular disorder; corneal infection;  
KW wound healing; epithelial cell proliferation; food additive.

XX  
OS Homo sapiens.  
XX  
PN MO200155312-AZ.  
XX  
PD 02-AUG-2001.  
XX  
PF 17-JAN-2001; 2001WO-US01321.  
XX  
PR 31-JAN-2000; 2000US-0179065.  
PR 04-FEB-2000; 2000US-0180628.  
PR 24-FEB-2000; 2000US-0184664.  
PR 02-MAR-2000; 2000US-0186350.  
PR 16-MAR-2000; 2000US-0189874.  
PR 17-MAR-2000; 2000US-0190076.  
PR 18-APR-2000; 2000US-0198123.  
PR 19-MAY-2000; 2000US-0205515.  
PR 07-JUN-2000; 2000US-0209467.  
PR 28-JUN-2000; 2000US-0214866.  
PR 30-JUN-2000; 2000US-0215135.  
PR 07-JUL-2000; 2000US-0216647.  
PR 07-JUL-2000; 2000US-0216880.  
PR 11-JUL-2000; 2000US-0217487.  
PR 11-JUL-2000; 2000US-0217496.  
PR 14-JUL-2000; 2000US-0218290.  
PR 26-JUL-2000; 2000US-0220963.  
PR 26-JUL-2000; 2000US-0220964.  
PR 14-AUG-2000; 2000US-0224518.  
PR 14-AUG-2000; 2000US-0224519.  
PR 14-AUG-2000; 2000US-0225213.  
PR 14-AUG-2000; 2000US-0225214.  
PR 14-AUG-2000; 2000US-0225266.  
PR 14-AUG-2000; 2000US-0225267.  
PR 14-AUG-2000; 2000US-0225268.  
PR 14-AUG-2000; 2000US-0225270.  
PR 14-AUG-2000; 2000US-0225447.  
PR 14-AUG-2000; 2000US-0225757.

PR 14-AUG-2000; 2000US-0225758.  
PR 14-AUG-2000; 2000US-0225759.  
PR 18-AUG-2000; 2000US-0226279.  
PR 22-AUG-2000; 2000US-0226681.  
PR 22-AUG-2000; 2000US-0226686.  
PR 22-AUG-2000; 2000US-0227182.  
PR 22-AUG-2000; 2000US-0227182.  
PR 30-AUG-2000; 2000US-0227009.  
PR 30-AUG-2000; 2000US-0228924.  
PR 01-SEP-2000; 2000US-0229287.  
PR 01-SEP-2000; 2000US-0229343.  
PR 01-SEP-2000; 2000US-0229344.  
PR 01-SEP-2000; 2000US-0229345.  
PR 05-SEP-2000; 2000US-0229509.  
PR 05-SEP-2000; 2000US-0229513.  
PR 06-SEP-2000; 2000US-0230437.  
PR 06-SEP-2000; 2000US-0230438.  
PR 08-SEP-2000; 2000US-0231242.  
PR 08-SEP-2000; 2000US-0231243.  
PR 08-SEP-2000; 2000US-0231244.  
PR 08-SEP-2000; 2000US-0231244.  
PR 08-SEP-2000; 2000US-0231413.  
PR 08-SEP-2000; 2000US-0233080.  
PR 08-SEP-2000; 2000US-0233081.  
PR 12-SEP-2000; 2000US-0231968.  
PR 14-SEP-2000; 2000US-0232397.  
PR 14-SEP-2000; 2000US-0232397.  
PR 14-SEP-2000; 2000US-0232399.  
PR 14-SEP-2000; 2000US-0232399.  
PR 14-SEP-2000; 2000US-0232400.  
PR 14-SEP-2000; 2000US-0232401.  
PR 14-SEP-2000; 2000US-0233063.  
PR 14-SEP-2000; 2000US-0233064.  
PR 14-SEP-2000; 2000US-0233065.  
PR 21-SEP-2000; 2000US-0234223.  
PR 21-SEP-2000; 2000US-0234274.  
PR 25-SEP-2000; 2000US-0234997.  
PR 25-SEP-2000; 2000US-0234998.  
PR 26-SEP-2000; 2000US-0235484.  
PR 27-SEP-2000; 2000US-0235834.  
PR 27-SEP-2000; 2000US-0235836.  
PR 29-SEP-2000; 2000US-0236327.  
PR 29-SEP-2000; 2000US-0236367.  
PR 29-SEP-2000; 2000US-0236368.  
PR 29-SEP-2000; 2000US-0236369.  
PR 29-SEP-2000; 2000US-0236370.  
PR 02-OCT-2000; 2000US-0236802.  
PR 02-OCT-2000; 2000US-0237037.  
PR 02-OCT-2000; 2000US-0237038.  
PR 02-OCT-2000; 2000US-0237039.  
PR 02-OCT-2000; 2000US-0237040.  
PR 13-OCT-2000; 2000US-0239935.  
PR 13-OCT-2000; 2000US-0239937.  
PR 20-OCT-2000; 2000US-0240960.  
PR 20-OCT-2000; 2000US-0241221.  
PR 20-OCT-2000; 2000US-0241785.  
PR 20-OCT-2000; 2000US-0241786.  
PR 20-OCT-2000; 2000US-0241787.  
PR 20-OCT-2000; 2000US-0241808.  
PR 20-OCT-2000; 2000US-0241809.  
PR 01-NOV-2000; 2000US-0244826.  
PR 01-NOV-2000; 2000US-0244617.  
PR 08-NOV-2000; 2000US-0246474.  
PR 08-NOV-2000; 2000US-0246475.  
PR 08-NOV-2000; 2000US-0246476.  
PR 08-NOV-2000; 2000US-0246477.  
PR 08-NOV-2000; 2000US-0246478.  
PR 08-NOV-2000; 2000US-0246523.  
PR 08-NOV-2000; 2000US-0246524.  
PR 08-NOV-2000; 2000US-0246525.  
PR 08-NOV-2000; 2000US-0246526.  
PR 08-NOV-2000; 2000US-0246527.  
PR 08-NOV-2000; 2000US-0246528.  
PR 08-NOV-2000; 2000US-0246532.  
PR 08-NOV-2000; 2000US-0246609.

PR 08-NOV-2000; 2000US-0246610.  
 PR 08-NOV-2000; 2000US-0246611.  
 PR 08-NOV-2000; 2000US-0246613.  
 PR 17-NOV-2000; 2000US-0249207.  
 PR 17-NOV-2000; 2000US-0249208.  
 PR 17-NOV-2000; 2000US-0249209.  
 PR 17-NOV-2000; 2000US-0249210.  
 PR 17-NOV-2000; 2000US-0249211.  
 PR 17-NOV-2000; 2000US-0249212.  
 PR 17-NOV-2000; 2000US-0249213.  
 PR 17-NOV-2000; 2000US-0249214.  
 PR 17-NOV-2000; 2000US-0249215.  
 PR 17-NOV-2000; 2000US-0249216.  
 PR 17-NOV-2000; 2000US-0249217.  
 PR 17-NOV-2000; 2000US-0249218.  
 PR 17-NOV-2000; 2000US-0249244.  
 PR 17-NOV-2000; 2000US-0249245.  
 PR 17-NOV-2000; 2000US-0249264.  
 PR 17-NOV-2000; 2000US-0249265.  
 PR 17-NOV-2000; 2000US-0249297.  
 PR 17-NOV-2000; 2000US-0249299.  
 PR 17-NOV-2000; 2000US-0249300.  
 PR 01-DEC-2000; 2000US-0250160.  
 PR 01-DEC-2000; 2000US-0250391.  
 PR 05-DEC-2000; 2000US-0251030.  
 PR 05-DEC-2000; 2000US-0251988.  
 PR 05-DEC-2000; 2000US-0256719.  
 PR 06-DEC-2000; 2000US-0251479.  
 PR 08-DEC-2000; 2000US-0251856.  
 PR 08-DEC-2000; 2000US-0251868.  
 PR 08-DEC-2000; 2000US-0251869.  
 PR 08-DEC-2000; 2000US-0251989.  
 PR 08-DEC-2000; 2000US-0251990.  
 PR 11-DEC-2000; 2000US-0254097.  
 PR 05-JAN-2001; 2001US-0259678.  
 (HUMA-) HUMAN GENOME SCI INC.  
 PA Rosen CA, Barash SC, Ruben SM;  
 PI  
 XX  
 XX  
 PI WPI: 2001-488782/53.  
 DR N-PSDB; AAS34287.  
 XX  
 XX  
 PT New polynucleotides and polypeptides for diagnosing, treating,  
 PT preventing or prognosing e.g. diseases or disorders of the nervous,  
 PT musculoskeletal, excretory, gastrointestinal, reproductive, and  
 PT respiratory systems -  
 XX  
 PS Claim 11; SEQ ID NO 1711; 642bp; English.  
 XX  
 CC The invention relates to novel nucleic acids encoding novel human foetal  
 CC antigens. The nucleic acids and proteins are used to prevent, treat (e.g.  
 CC by gene therapy) or ameliorate a medical condition in e.g. humans, mice,  
 CC rabbits, goats, horses, cats, dogs, chickens or sheep. They  
 CC are also used in diagnosing a pathological condition or susceptibility  
 CC to a pathological condition. The antibodies to the antigens can also  
 CC be used in alleviating symptoms associated with the disorders and in  
 CC diagnostic immunoassays e.g. radioimmunoassays or enzyme linked  
 CC immunoassay assays (ELISA). Disorders which are diagnosed or treated  
 CC include autoimmune diseases e.g. rheumatoid arthritis,  
 CC hyperproliferative disorders e.g. neoplasms of the breast or liver,  
 CC cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders  
 CC e.g. cerebral ischaemia, angiodenesis, nervous system disorders e.g.  
 CC Alzheimer's disease, infections caused by bacteria, viruses and fungi  
 CC and ocular disorders e.g. corneal infection. The polypeptides can also  
 CC be used to aid wound healing and epithelial cell proliferation, to  
 CC prevent skin aging due to sunburn, to maintain organs before  
 CC transplantation, for supporting cell culture of primary tissues, to  
 CC regenerate tissues and in chemotaxis. The polypeptides can also be used  
 CC as a food additive or preservative to increase or decrease storage  
 CC capabilities, fat content, lipid, protein, carbohydrate, vitamins,  
 CC minerals, cofactors and other nutritional components. Numerous  
 CC examples of diseases and disorders treated by the nucleic acids and

CC proteins are given in the specification. The present sequence

Alignment Scores:

Pred. No.:	1.25e-40	Length:	79
Score:	49.00	Matches:	49
Percent Similarity:	100.00%	Conservative:	0
Best Local Similarity:	100.00%	Mismatches:	0
Query Match:	9.68%	Indels:	0
DB:	22	Gaps:	0

US-09-745-506-74 (1-1553) x AAU21467 (1-79)

OY 596 CTTGAGCTGTACCTCCAGCCATCATCTTCCAAAGCTCCCAACTACCTACAG 655  
 |||||  
 DB 15 LeuGlyAlaCysThrSerAqProIleHisProSerIysAlaProAsnIYrProThcIu 34  
 |||||  
 OY 656 GGAACCCAGCAGTAGAATTCAACGTTAACTACACCCAGACCTGGACAAAGTCATGCT 715  
 |||||  
 DB 35 GlyAsnHisArgValGluPheAsnValAsnIYrThrGlnAspLeuAspIysValMetSer 54  
 |||||  
 OY 716 GCAGTGAAGAAATTCAGCGCTTCT 742  
 |||||  
 DB 55 AlaValIysGlyIleAspGlyValSer 63  
 |||||

RESULT 15  
 AAU27916  
 ID AAU27916 standard; Protein: 146 AA.  
 AC AAU27916;  
 XX  
 XX  
 DT 18-DEC-2001 (first entry)  
 XX  
 DE Human contig polypeptide sequence #69.  
 XX  
 XX Mammal; human; rhesus monkey; baker's yeast; fission yeast; Norway rat;  
 XX mouse; Chinese hamster; African clawed frog; fruit fly; dog; leukemia;  
 XX cancer; lymphoma; neuroblastoma; autoimmune disorder; cell proliferation;  
 XX nervous system disorder; inflammatory disorder; cell differentiation; burn;  
 XX angiogenesis; stem cell growth factor; activin; inhibin; cartilage; bone;  
 XX genetic disorder; bone regeneration; tendon; ligament; tissue repair;  
 XX cytoskeletal; anti-rheumatic; antiarthritic; vulnery; antiinflammatory;  
 XX antibacterial; immunosuppressive; vasotropic; antiparkinsonian;  
 XX neuroprotective; osteopathic; antidiabetic; antistimatic; antiallergic;  
 XX immunostimulant; analgesic; gene therapy.  
 OS Homo sapiens.  
 OS Synthetic.  
 OS  
 PN WC200164834-A2.  
 XX  
 XX  
 PD 07-SEP-2001.  
 XX  
 XX 26-FEB-2001; 2001WO-US04926.  
 XX  
 XX 28-FEB-2000; 2000US-0515126.  
 XX 18-MAY-2000; 2000US-0577409.  
 XX 17-JUN-2000; 2000US-0597707.  
 XX 14-JUL-2000; 2000US-0616807.  
 XX 19-SEP-2000; 2000US-0664641.  
 XX  
 XX (HYSE-) HYSD INC.  
 PA  
 XX  
 XX Tang YT, Liu C, Zhou P, Asundi V, Zhang J, Zhao QA, Ren F;  
 PI Xue AJ, Yang Y, Wehrman T, Wang J, Ma Y, Wang D, Chen R, Xu C;  
 PI Dmanac R;  
 XX  
 XX WPI: 2001-589862/66.  
 DR N-PSDB; AAS44816.  
 XX  
 PT Novel polypeptides and nucleic acids obtained from cDNA libraries  
 PT prepared from various human tissues, for diagnosis, treatment of  
 PT cancer, neurological, inflammatory disorders and for use in arrays for  
 PT detection -

XX Claim 10; Page 132; 153pp; English.

XX  
PS Sequences AAU27676-AAU28019 represent full-length polypeptides and  
CC contig polypeptides of the invention. The proteins and their associated  
CC DNA sequences are useful for the treatment, diagnosis and prevention of  
CC various types of disorder in a mammalian subject such as a human, dog,  
CC monkey, mouse, hamster or rat. The disorders include cancers such as  
CC leukemia, lymphoma and neuroblastoma, autoimmune disorders such as  
CC multiple sclerosis, connective tissue disease, rheumatoid arthritis,  
CC diabetes mellitus, allergic rhinitis, asthma and eczema, nervous system  
CC disorders such as Parkinson's disease, Alzheimer's disease, Huntington's  
CC chorea, amyotrophic lateral sclerosis, spinal muscular atrophy and  
CC Wernicke disease, inflammatory disorders such as nephritis, Crohn's  
CC disease, ischaemia-reperfusion injury, shock, sepsis and inflammatory  
CC bowel disease. The sequences exhibit activity relating to angiogenesis,  
CC cell proliferation, cell differentiation, stem cell growth factor,  
CC activin or inhibin. Therefore, they can be used to manipulate stem cells  
CC in culture to give rise to neuroepithelial cells that can be used to  
CC augment or replace cells damaged by illness, accidental damage or genetic  
CC disorders. The sequences may also be used for regeneration of bone,  
CC cartilage, tendons and ligaments and in tissue repair and burn healing.  
CC Note: Some sequences for this patent did not form part of the printed  
CC specification, but were obtained in electronic format directly from WIPO  
CC at ftp.wipo.int/pub/published\_pct\_sequences.

XX  
SQ Sequence 146 AA:

Alignment Scores:  
Pred. No.: 1.5e-34 Length: 146  
Score: 43.00 Matches: 43  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 8.50% Indels: 0  
DB: 22 Gaps: 0

US-09-745-506-74 (1-1553) x AAU27916 (1-146)

OY 257 GCTCTCCTTTCCTTGAATGACTTTCATCCCTGCTGTCGAGAGTTGGACAAT 316  
|||  
DB 36 AlaIleuSerSerIleuAsnAspPheAlaSerIleuSerPheAlaGluSerTrpAspAsn 55  
OY 317 GTTGATTACTGCTGAGACCAAGCCACACATACTGTAATACACTCTTCTGACAAT 376  
|||  
DB 56 ValGlyIleuLeuValGluProSerProHisThrValAsnThrIleuPheIleuThrAsn 75  
OY 377 GACCTGACT 385  
|||  
DB 76 AspIleuThr 78

Search completed: August 22, 2003, 14:32:24  
Job time : 112 secs